

Amendments to the Specification:

Please insert the following paragraphs at page 34, line 5 after

“(http://www.sanger.ac.uk/DataSearch/gq_search.shtml) (Table 2)” and before “As shown in Table 2, ...” as with the following amended paragraph:

-- The sequence of the V_H region of the DP47 germ line is as follows (SEQ ID Nos:113 and 114):

1	2	3	4	5	6	7	8	9	10	11	12	13	14	15
E	V	Q	L	L	E	S	G	G	G	L	V	Q	P	G
gag	gtg	cag	ctg	ttg	gag	tct	ggg	gga	ggc	ttg	gta	cag	cct	ggg
16	17	18	19	20	21	22	23	24	25	26	27	28	29	30
G	S	L	R	L	S	C	A	A	S	G	F	T	F	S
ggg	tcc	ctg	aga	ctc	tcc	tgt	gca	gcc	tct	gga	ttc	acc	ttt	agc
31	32	33	34	35	36	37	38	39	40	41	42	43	44	45
S	Y	A	M	S	W	V	R	Q	A	P	G	K	G	L
agc	tat	gcc	atg	agc	tgg	gtc	cgc	cag	gct	cca	ggg	aag	ggg	ctg
46	47	48	49	50	51	52	53	54	55	56	57	58	59	60
E	W	V	S	A	I	S	G	S	G	G	S	T	Y	Y
gag	tgg	gtc	tca	gct	att	agt	ggt	agt	ggt	ggt	agc	aca	tac	tac
61	62	63	64	65	66	67	68	69	70	71	72	73	74	75
A	D	S	V	K	G	R	F	T	I	S	R	D	N	S
gca	gac	tcc	gtg	aag	ggc	cgg	ttc	acc	atc	tcc	aga	gac	aat	tcc
76	77	78	79	80	81	82	83	84	85	86	87	88	89	90
K	N	T	L	Y	L	Q	M	N	S	L	R	A	E	D
aag	aac	acg	ctg	tat	ctg	caa	atg	aac	agc	ctg	aga	gcc	gag	gac
91	92	93	94	95	96	97	98	99						
T	A	V	Y	Y	C	A	K	.						
acg	gcc	gta	tat	tac	tgt	gcg	aaa	ga						

The sequence of the VL region from the DPK15 germ line is as follows (SEQ ID Nos:115 and 116):

1	2	3	4	5	6	7	8	9	10	11	12	13	14	15
D	I	V	M	T	Q	S	P	L	S	L	P	V	T	P
GAT	ATT	GTG	ATG	ACT	CAG	TCT	CCA	CTC	TCC	CTG	CCC	GTC	ACC	CCT

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16	17	18	19	20	21	22	23	24	25	26	27	28	29	30
G	E	P	A	S	I	S	C	R	S	S	Q	S	L	L
GGA	GAG	CCG	GCC	TCC	ATC	TCC	TGC	AGG	TCT	AGT	CAG	AGC	CTC	CTG
31	32	33	34	35	36	37	38	39	40	41	42	43	44	45
H	S	N	G	Y	N	Y	L	D	W	Y	L	Q	K	P
CAT	AGT	AAT	GGA	TAC	AAC	TAT	TTG	GAT	TGG	TAC	CTG	CAG	AAG	CCA
46	47	48	49	50	51	52	53	54	55	56	57	58	59	60
G	Q	S	P	Q	L	L	I	Y	L	G	S	N	R	A
GGG	CAG	TCT	CCA	CAG	CTC	CTG	ATC	TAT	TTG	GGT	TCT	AAT	CGG	GCC
61	62	63	64	65	66	67	68	69	70	71	72	73	74	75
S	G	V	P	D	R	F	S	G	S	G	S	G	T	D
TCC	GGG	GTC	CCT	GAC	AGG	TTC	AGT	GGC	AGT	GGA	TCA	GGC	ACA	GAT
76	77	78	79	80	81	82	83	84	85	86	87	88	89	90
F	T	L	K	I	S	R	V	E	A	E	D	V	G	V
TTT	ACA	CTG	AAA	ATC	AGC	AGA	GTG	GAG	GCT	GAG	GAT	GTT	GGG	GTT
91	92	93	94	95	96	97	98	99	100	101				
Y	Y	C	M	Q	A	L	Q	T	P	P				
TAT	TAC	TGC	ATG	CAA	GCT	CTA	CAA	ACT	CCT	CCT	---			